

OIPE

## RAW SEQUENCE LISTING

DATE: 01/29/2003 TIME: 15:14:26 PATENT APPLICATION: US/09/830,913A

Input Set : A:\pb0005usn seqlist.txt Output Set: N:\CRF4\01292003\I830913A.raw

```
4 <110 - APPLICANT: INCYTE PHARMACEUTICALS, INC.
              WALKER, Michael G.
              VolkMUTH, Wayne
              KLINGLER, Tod M.
      3 -(10) - TITLE OF INVENTION: CORTICOSTEROID SYNTHESIS-ASSOCIATED GENES
     11 -: 130 - File REPERENCE: PB-0005 PCT
C--> 13 <140> CURRENT APPLICATION NUMBER: US/09/830,913A
C--> 14 <141> CURRENT FILING DATE: 2003-01-29
     1. Histor Paide Application Number: 60/155,269
     17 -: 151 - PRICE FILING DATE: 1998-11-06
     1 4 - 1140 - NUMBER OF SEQ ID NOS: 9
     21 - 2170 | SOFIWARE: PERL Program
     .4 +211 + LENSTH: 3736
             TYPE: DNA
     16 -1113 - 05.GANISM: Homo sapiens
     LA FULLO FEATURE:
     . * HOLL: MAME: KEY: misc_feature
     -4 - 223 - DIHER INFORMATION: Indyte ID No.: 064973CB1
     F1 -1400 SEQUENCE: 1
        -totitopagt tugagaagag gagtiggtigat aaagoobact gacottoaca battotaaaa -60\,
        laitygttygo satyccagta taccaabaat batgatbago tysaagaaab agatatttta 120
         ustigopaga abaabaaatg atgaggbaab tabagtbaga tetatagoba gobatotato 180
     50^\circ aponotagas gottopagag abagtgaaab tgbaagatgo totbaabagg attatgtoto 240^\circ
         atggagadda gtaagaaaat batttatotg aaggtgaaat gbagagttgg ataagaaata 300
     38 - osthgotogg tytotaaaat gotogoottoo tooototaot obacotooat bootogaott 300
     50 -thysocotty ghotagyago otaagyabot toacocotyt ycaccacca ayaaagayga 4.\%
     40 -alabettiggs throsecttig galaxtyotgy ggtbootggt gtggtaagaa actoaabato 4\%
        -amabgggta: ghagaaggat gitottotgg gattigoagg tabataaaaa atgtatggba 540
     40^\circ touthbooth ghasabboth coaghtbook agrigagaagy ggagbaggtg bbbactgaty 6\%
         \eta_{	ext{Ada}}agytat gutgotatgt tyatqtgtaa qtqaaatcay tiqigtgcaa tagacagygy 660
     - oututtoaty ggagoatoag obagittota sasobbabay gobatbagba gotagaggity 7.0
     .; .;
         yet qotttig godagadatg gaddotaaat daadagadaa tiggdattigto gaagagdaad 7\%
         otyttaatga atpatyttaa aaatbaaggt ttygettoay titaaatbab ttyaggtaty 841
     40
        -algorithator typicitopag agatalaabat alagorigatot toocaalaata obatcattag 9\%
     .; 7
         quaditations assaulted tagittettt egittgettg tettetgett tgtttettgg 900
     .; :-
     44 -beaugopath cappabagab thotogopological agotogagaga balatogotogotalataa 1/20
         gyatottiga thaabbooda taaggoatgi yigigiatad aaatatabii otottiggoi 1780
         introgadata gwadotoago tigttaadoaa jigggaaatao atbagatotig baababagaa 1140
         atgritotiges transititios appatigosta gigasticasop patititatosa gigtottibity 1.\,30
        -gathtyttta atoaataago ootataatoa ottyotaaab astyygotto atoaccbagg 1.\,60
     53
         gataaaaaca gigatbattg tottggadot dotgdatday ootattdaaa attatototd 1520
         tototagott tonacaaato otaaaattoo tgtoocaago caccoaaatt otoagatott 1380
```



.1/2#.2000 TINE: 18:14:1 FATENT APPLICATION: US/09/830,913A

Input Set : A:\pb0005usn seqlist.txt Cutput Set: N:\CRF4\01292003\I830913A.raw

```
ttotggaada aggoagaata taaaataaat atacatttag tggottgggo tatggtotoo 1440
        aaagatooti caaaaataca toaagooago ticaticact caciitacii agaacagaga 1800
       tataagggoo tgggatgoat ttattttato aataobaatt tttgtgggooa tggoagacat 1560
59 tgotaateaa teasagoast atttoetatt aageesaetg atttetteas aatesttete 1620
50^{\circ} aaattacaat toossaagago ogobactosa bagtosgatg aaccossabag toagatgaga 1680^{\circ}
\#1 -gamatqaaco otaottigota tototatott agamagomma aacamadagg magtttocagg 174\%
v2 -qagaatggga aaqobagggg goataaaagg tabagtbagg ggaaaataga totaggoaga 1600
63. qtgcottagt baqqqabbac gggogdtgaa totgbaqtgo baababbaaa btgababato 1860
64^{\circ} throaggigta potopaappo tagopticip poacagoigo piacaacaga giptopocago 1920
        oftotoagag agotaaaaoo agaaatttoo agactoatga aagcaacooo coagoctoto 1400
11.1
        \phi meansoring engentiaged that that an analytic and the states of a typical states and the states of the states 
1.
         thataagbag gggbbagaat abobbagbba bobgbagtga babbgbtgga bobbtgaaaa 1000
        \odotpathopata ggagaatggg thoobboaggo hoabagtgta gagabattga gobbatbaba 1101
1
\mathcal{C}^{+} autiquiting objections totalaalaga totaloosed coatgreat graphysis \mathbb{R}^{1}
^{600} -thoograpse attoagaagt toaagbogag atgotgacgt tgotgagbaa ogagatggtg _{\odot} \%
The agree to agree of the agree
         quegettogg casadoatet tgatggasts gggasetges satgtstgat gatettgass 2400
        Aggotbagoa ggattigtto tiaaadbgab tbagtgtgto atbobbggtt atttagaatt 1400
74 ahagttaada addaqaaadt totataagad tgtatgaada aggtgatato ttoatagtgg 1910
       gotattadag gdaggaaaat gttttaabtg gtttadasaa tooatdaata ottgtgtbat 1950
^{\circ}M, topotytasa agypagyaga batytyatta tyatbayyaa abtybabaaa attattytti.^{\circ}M)
       -toageocoog Equiatogic offitigaact giftiffit fitaliaaage caaattigig arphi
78 engbatatab togtattoda tgtgttagat ggaagdattt ootatodagt gtgaataaaa . 😘
99 Agaadagttg tagtaaatta ttataaagoo gatgatattt batggoaggt tattotabba 1920
80 -agotytyctt gttgyttitt oodatgadty taitgoittt ataaatytad aaatayttad . 88\%
       Agaaatgaog agaooottgt tigbabagba tiaataagaa bottgataag aabbatatto 194\%
         tyttyadago dagotdabag titottydot gaagottygt ybabbotbba ytgagababa 2000
13
         aratototot titaboaaag tigagaadag agotggigga tiaattaata gibtibgata 2002
64 thtggodatg ggtaadotoa ttgtaadtat batbagaatg ggdagagatg atbttgaagt 11110
       Apidudatada obaaagtoda aadadtatgt dagatggggg taaaatodat taaagaadag (1960
        quannastas tratasgatg atasgosast gtttpsgppp aatgtpsaco pagttasass NAC
A/ Maaaattaat gotgtgtaaa atggttgaat tagtttgcaa actatataaa gacatatgca FEW
         gtaaaaagto tgitaatgoa catooigigg gaatggagig tootaaccaa tigcotitto 1960
44 tightatotg agototoota tattatoata otoagataao baaattaaaa gaattagaat 1426
90 andarittita atabactical cattagacto tictagetti etietittitg tyataatica 34%0
AI -quadrataget abggatoses autgeototy agreattyte abadadaats agetatodot ^{10}\,M_\odot
        ataboatgot ataggagaot gggbaaaaob tgtabaatga baabbotgga agttgbtttt 3600
93 -titasaassa taataaatti ottaastoss otottittibo tyyttytoty tityttätää ^{4}e60
(4) automasort attoaagtod toaatatoot gatoataata doatgotata ggagaotggg 37.00
       -baadabbtgt adaatgadaa obotggaagt tgottottta aaaaaaataa taaatttott 9780
                                                                                                                                                                  3736
196 - Asatoaaaaa aaaaaa
93 <210 - SEQ ID NO: 2
100 -0011 - LENGTH: 1187
1:1 -L.1L - TYPE: DNA
1/2 -L1: ORGANISM: Homo sapiens
1.4 -0.20 - FEATURE:
198 -0.21 - NAME/KEY: misc feature
196 - M21 - OTHER INFORMATION: Incyte ID No.: 965781CB1
108 <450 - SEQUENCE: 2
```

## RAW SEQUENCE LISTING

FATENT AFFLICATION: US/09/830,913A

Input Set : A:\pb0005usn seqlist.txt Output Set: N:\CRF4\01292003\I830913A.raw

CATE:

TIME: 18:14:5

```
atatacgatt actataggga titiggcopic gaggcaagaa attoggcacg aggatcagac 🕬
          cateagaagg attigiataa agagigasis isotaigaag giaaaggesa esestetisa 120
          guidoaguga orgagatada utulibedaat oolgggggga aatadagada dagdaagtid 180
         -cttcttccct ttggaaattt ggcagotgoc ttcaccagtg agcacaaago cacatttcaa 240
     113 aggaaastga daaattatos deagstgoda gaagaagaaa tootoaetgg asggottost.
         gittootgig giloattato igatiggoig bagggatgaa agittitaag tibalaggab 360
         tgatgatict cotcacciet gogitticag coggiticagy acaaagtoca atgactytyc 420
     116 tgtgeterat agaetggtte atggteaeag tgeaebeett batgetaaab aaegatgtgt 480
     11..
         gtqtacactt toatgaacta bacttqqqoo tqqqttqcoo bobaaacbat qttcaqocac 540
         Lubgostarda gittoapotad ogigitaditg aatgiggdat bagggodaaa goigitoidid 600
     11:
     119
          -aggraphized tatotabago abigagatab abiaototio taagggbabg bbatotaagt 640
     1.3
          ittytyatood agtytoatyt yotyodooda aaaaytoodd atyyotdadd aagdootydt 77.6\,
         poatgagagt agodagbaag agbagggoda bagbobagaa ggatgagaaa tgotabgagg 781
     1. 1
          tyttoagott gibabagtoo agtbaalaggo bbaabtybga ttytbbabbt tytytottba 841
         ytgaagaaga goataoobag gtooottgto accaagbagg ggotbaggag gotbaabotb 900
     1 . .
        -tgcapocato beactitett gabatitety aggatiggto tettoacada gatgatatga 90\%
     1. 4
     108 Thiggground greateston ggottggggt etestgaaga tgetatttet agaattagta 1920
         - tatagtigtas awatigtistiga isaaataagtig otisttigtigas ootisatigtiga goastittiga 1.80
     1....
          gaaagagaaa ootatagdaa ootoatgaat taagoottet totatattet tatattoatg 1140
     1.
         tgthaabaaa aaataasata saattotgat ogbataasaa aaasaaa
                                                                               11.7
     1.
     1:1 :::110 - SEQ ID NO: 3
     1:. KU11 - LENGTH: 1395
     199 - Miller TYPE: DNA
     1:4 -0013 - ORGANISM: Homo sapiens
     130 KINGO - FEATURE:
W--> 137 <221> NAME/KEY: misc feature
     136 HO23 - OTHER INFORMATION: Incyte ID No.: 1419725081
     140 H400 - SEQUENCE: 3
          otokaaggda aacaaaagga aatgoocggo tooocatggo tytyyocayo acottoatac -6\%
     14:
         , usquigotoss obotidagaso potoattista tipodagigota agastoopas titagogotoo 1,\, \emptyset
         -negetigety olyggagtag gggtalaggg liggaggaatt aagetyetta gagealyggg 1 e \delta
     14:
     144 -beaugeggae aggitiggibt gggaggetga ggibtobatg gitggtggatg gaaaaggaga 440
     14% Aggsagggad tagtgagada dogggtotgo obcaaagott tgtgttgtto totopatggg 300
     146
         laabagggtaa bagbagagga aggatogaaa goobaggbag agaabttaga aatgobtgot 360
         -boasatatat ababtgotga gtobagaaag atbagggott bagaggtotb bybttotgbb 410\,
     14/
         totogggatg goggggdagg gogattotogg tgagtgggag agtototogot goaggtabad 480
     14.
     149 tggabactgo obactactto ggttoagogt gggobagado tatgggobagg tgabtggtba 840
```

-fotworthags ggoddtootg godtagootg goddootgto dadogdadad thotgodtod  $600\,$ 

-pattoggoot omaagatoto otgaggttoo bagggagagt blabotgtba ggogtgggba 660

Agahaggoto agotodagoa tgatodotgg gtacacaggt ttgtacgcag gtatgcacag  $720\,$  itgusested saggageaty tyttodaaad adtaasgagt ettesttyte setgestyde 780 184 -casultetigt accordages cagiticated tigopaagaa cigoagooag gibigggoog 640

-aggetotigay tigabittabi babitigbatig saaagbaagg gagtigaagag biabbaaagg  $9\omega$ 

agmicaagmg aaraaagmad acagagaaamg accaggtgoc agagcoggag gggcagctgg 960

Aggagoogad actggaggtg gtggaadaag ottotoocta otocatggat gadagggado 1000 185 oboggaagit ottoatgica ggdactgdag gaattigggd agaagdacto accaggdagt  $1.60\,$ 

goodaggado odaaabatot oodoodaatti oodagaadat abootoagaa ootgggibti 1140 -thaddtaadt atgggggdta dytgddaggg tataagttod agtttggdda dadatttggd  $1000\,$ 

catetoacco atgatgetot gggoeteage acetteeaga agoagotett ggettagged 1200

1:0 1°:

1°.

154

1: +

1:7

1:.4

100

161

## RAW SEQUENCE LISTING

FATENT APPLICATION: US/09/830,913A

Input Set : A:\pb0005usn\_seqlist.txt
Output Set: N:\CRF4\01292003\I830913A.raw

TATE:

```
-adiggarato aagitoodit oddittidat oddatoodag odatroitit ggaagggaga 1820
         - gaggtgggtg ggagggtggg ggaacacaaa gagaaaatgq tittggaggct gagcaccitt 1390
     164 - titattaata ggiat
     167 <2105 SEQ ID NO: 4
     168 K211 - LENGTH: 36°
     169 K212 - TYPE: DNA
      170 K213 - ORGANISM: Homo sapiens
     172 < 220 + FEATURE:
W--> 173 <221> NAME/KEY: misc feature
     174 KRRE OTHER INFORMATION: Incyte ID No.: 23645820B1
     176 -:401 - SEQUENCE: 4
     177
          stittetbag baaagygeag eactgetege tittgetgggg eagebetbeb aggaagateb - 60
     17.
          -bagaagaat gototgaggt gigigaabab igigaatgib atggabaaag bibagogigg 1/\phi
     17\% gamuscatgy gasticaggs obtgaaagsa goagatotoa obtgoattgy aggatgagas 1\%
     18% genecatgaa gwagaaagaa gabgobabag abagbagbagbab agtabbabbt tobagagata 24%
     181 grateactyt gadaestagt teggaatyca atggegogat ettygetaac tyaaaootee 3.0
     18. quitospagg thosasagst totottgoot dagosttoog syttygg sotsosgsas 860
     18: acceptotate aggreeocca gatagitatea gasagetgas gatticoaga tegetgoate 41\%
     184 Intercaphas appropriates operquetica actacotact gootgetigag casotoctac 440
     18) Attakantost auctoratts gotasagaga pagattotag aptigotop satatosigg (40)
                                                                                 5.67
     18) Stypatbabb agbaataaag bottabt
     189 - 0110 - SEQ ID NO: 5
     190 HILL - LENGTH: 920
     191 A. L. TYPE: DNA
     192 Hull: ORGANISM: Homo sapiens
     194 - C. FOR FEATURE:
W--> 195 <221> NAME/KEY: misc feature
     196 HIJB - OTHER INFORMATION: Incyte ID No.: 2737624081
     19: K400 - SEQUENCE: 5
     199 -gogmanacot ghagtegtag otabitggga ggotgaggda ggggaatbab tigaabbbbg \sim 60
     26c -wag roggagg thytagtggg cogagattgt godabtgdg todagogtgg gogadagagt 17\%
         - daduoticcat oticaalaaaa aasaaaaaaa aatotatigot agtagattad aactticadad 190
     \mathbb{R}^{0} . Tagaggagtt otiggadaaag ottttaatta gtoaaadtaa attaaggotd attaaaagga \mathbb{R}^{4} \mathbb{R}
     30\% haggaactas tiggaaatta tigdaattoaa taatttagad totigttadoa gigatotttoa 30\%
     \pm 0.04 taasaasttta atttopataa toataabota aatgagttot taaaagaatto tataagbaat 3\%
     20) – agotgattaa toogoobotgo aagatgaaga ttataabtgo boatttacoo aattaaaago 420
     \pm 900 -alaggoagby chalaitatga gaggataaad aatattagtt aacatttotg ttatttatga 480
     N(\mathbb{N}) typosaattay tagtaayata attocacago tytosaotti yttiggyyst yydaacttoi fM(\mathbb{N})
         intghetakan ampotkasag titagtatio igggagaagi ggoiggaaga aggggiaata 🕕 🤄
     LCF togetgaaago awattooott toodaggagt daagagaatt tatgtgaggt ggotoodgod \kappa 60
     210 tgtwatetba gesetttegg aggecaagge aggeaatbac etgagggeag gagaccagec 700
     LII tgaqtaaqat gqsgasacco qatqtqtaqt aaaaataqaa aatgagqtgg gogtggtggt %60
         - qoangootgt awtoobagot acttgggagg otgaggbagg agaatogoot gaabooggga 840
         - ogongaqitt gasqtiaagob aagatojago cactgtacto cagootgggg gacagatoga ^{140}C
                                                                                 920
         - agotgtotta aasaasaaaa
     117 -0110- SEQ ID NO: 6
     L18 -07110- LENGTH: 1564
     21" KU130 TYPE: DNA
```

220 KL139 ORGANISM: Homo sapiens

## RAW SEQUENCE LISTING

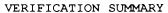
FATENT AFELL MATERN: US/09/830,913A

: ATE:

1 .... TIME: 19:14:.

Input Set : A:\pb0005usn seqlist.txt Currut Set: N:\CRF4\01292003\I830913A.raw

```
222 KULD FEATURE:
203 - 2010 NAMES KEY: misc insture
224 - 223: OTHER INFORMATION: Indyte ID No.: 28670650B1
226 kilolu SEQUENCE: 6
     labritti 44 aast rotte tilaasigit tosigattat scatigilig casagtaasi
    i gaaloaggitti aannthotyg argottanit ittaghagaaa tatathagat aghaaacttig 12
    i imgubagang aan nabagot, titgoolianot, ghaatgagda tabali gittgi galibobagan Ibi
    -totattiong gognorgoan acgeaagnee coaactegen gaggagaget gincagenat 240
    ggorgotyto acadantaes tytatotity coagnitage tygatysica tipagtotyt 300
     gaarttotgg taogtgotgg tgatgaatga tgagdabada gagaggogat atdtgotgtt 360
    ttteettetg agitggggae taccagettt tgtggtgatt etecteatag ttattttgaa 420
US4 aggmatotat patpagagoa tytpadagat otatygapto attdatygty addtytyttt 480
135 tathocasad grotatgorg offigitoac typeagefort grideritga egigeologi 540
    ggtagtgtto gtggtyttba tobatqoota bbaggtgaag bbacagtgga aagcatatga 600
13€
    tgargtotic agaggaagga caaatgotgo agaaattoca otgattitat atototitgo 660
   - totuattioo gigadaiggo tiiggggagg actacadaig gootacagad acticiggai 720
.339 gut/gitloto titgipatti toaabagibt goagggabti taigittiba iggittatti 780
     batittadad aadbaaatgt gttgoddtat gaaggobagt tabactgtgg aaatgaatgg 84\%
\pm41 -geatostyga pocagoadag potititicad geologygagt gyaatgooto etgetyyagg 900
    - igawatcago aagtobacco agaatotoat oggtgotatg gaggaggtgo cabotgabtg 960
    . Igadagagba toottobaab agggbagtba ggbbagbbbt gatttaaagb baagtbbaba 1820
1.44 -raatiggaged abgotobogt bototiggagg atlatiggobag gggtdabtga talgoogatga 1.780
    - įgagtoccag gagttigatg atttaatatt tgčattaaaa actggtgotg gtotcagtgt 1140
    -bagigataat gaatotyyto aayydagoda yyayyyygo addttyadty adtoodayat 1200\,
\underline{L} \in \mathcal{C}_{k}
    -typogagoto aggagyatas obatogooga bactbabotg tagbabotba otaaccattb 12\,60
347
     yabrigaqbad abbbbbatab bogbatbago bottgobata aaabbbbbta agbadabbba 1320
1 1
    ostytytaat aggaasstyt gaattytast ygatyattaa tasaaasyty attyttytät 1340
     htgdagtata aattabtgat tgtatgtgad otgaaaatto abtgotataa gaaaggtgga 1440
    - publigitigi alwagitaat aggatgitoa tattocaagg atattagitig tittittaat 1870
     Batubbatat gyotaabatt giitaatgaa agiaataato aataaagbaa tagaasotaa 1600
    1934
USE <210. SEQ ID NO: 7
MET <211 - LENGTH: 752
MER KRIZH TYPE: DNA
∴5.2 <∴13 + ORGANISM: Homo sapiens</p>
381 KB23 - FEATURE:
202 K221 MAME/KEY: mist feature
263 K223 FOTHER INFORMATION: Incyte ID No.: 2961563CB1
265 <400. SEQUENCE: 7
\pm 66 gatqgqgqgqgqggaqg atqaaaaaat baccattacc abbtqaqbt titbcattot = 60
2600 light-thaasa goattyatga tittgoaccat titggagatag agototygot tattitattat 13\,
\pm 68 -aaa_{
m c}aaggaaqaa ooggaabag tiggtigootad ooacaatigoo agotaactotig aaggatigagg 14
241 tgggagaatt gotggagggo aattogaaac bagtotgggo aagatagtga gabotbatot 241
     caaaaaaaaa aaaagaagga aaaaacaaag gagttgatga aaccaaatta tgacaaacta 3.1
270
    aattaaaaaa tiaaaaaata goattagtgg gaacactgtg gtatttootg aggtattaaa 361
   -attitiacaac tagtaaaggi attiattata attiattogg attifttott agittattit 4\mathbb{I}
272
    tgacatggat aaaattattg titatotagg agootcagag tigagagagg atagtigtag 45
274
    agaagtatti otaaaatyga attatoatoo actaccagot aagagaaaat agitactigi 54)
275 ttaaagatab ottaaagbaa datatattgt agtitbaaga gaabaggotg abagtbaaga 600
```



HATENT ARRIGITATI N: US/09/830,913A

Imput Set : A:\pb0005usn\_seqlist.txt

Surrut Set: N:\CRF4\01292003\1830913A.raw

L:18 M:271 O: Current Application Number ditters, Replaced Current Application Number 1:14 M:271 O: Current Filing Date ditters, Replaced Current Filing Date L:137 M:257 W: Feature value mis-spelled or invalid, <221> Name/Koy for SEQ ID#:3 L:173 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:4 L:195 M:257 W: Feature value mis-spelled or invalid, <221 Name/Key for SEQ ID#:5